RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/712,359A
Source: IFW0
Date Processed by STIC: 10-25-04

ENTERED

IFWO

RAW SEQUENCE LISTING DATE: 10/25/2004
PATENT APPLICATION: **US/10/712,359A** TIME: 11:45:50

Input Set : **A:\27908007.app**
Output Set: **N:\CRF4\10252004\J712359A.raw**

<110> APPLICANT: CHANG, Y-H
VETRO, J.A.
MICKA, W.S.
<120> TITLE OF INVENTION: DOMINANT NEGATIVE VARIANTS OF METHIONINE AMINOPEPTIDASE 2 ("METAP2") AND CLINICAL USES THEREFOR
<130> FILE REFERENCE: 2790/66153/8007
<140> CURRENT APPLICATION NUMBER: 10/712,359A
<141> CURRENT FILING DATE: 2003-11-13
<150> PRIOR APPLICATION NUMBER: 09/943,123
<151> PRIOR FILING DATE: 2001-08-30
<160> NUMBER OF SEQ ID NOS: 26
<170> SOFTWARE: PatentIn Ver. 3.2
<210> SEQ ID NO: 1
<211> LENGTH: 71
<212> TYPE: PRT
<213> ORGANISM: Homo Sapiens
<220> FEATURE:
<223> OTHER INFORMATION: Human polylysine
<400> SEQUENCE: 1

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Lys Lys Lys Arg Arg Lys Lys Lys Lys Ser Lys Gly Pro Ser Ala Ala
 1               5                  10                  15
Gly Glu Gln Glu Pro Asp Lys Glu Ser Gly Ala Ser Val Asp Glu Val
            20                  25                  30
Ala Arg Gln Leu Glu Arg Ser Ala Leu Glu Asp Lys Glu Arg Asp Glu
        35                  40                  45
Asp Asp Glu Asp Gly Asp Gly Asp Gly Asp Gly Ala Thr Gly Lys Lys
    50                  55                  60
Lys Lys Lys Lys Lys Lys Lys
65                  70
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<210> SEQ ID NO: 2
<211> LENGTH: 71
<212> TYPE: PRT
<213> ORGANISM: Mus musculus
<220> FEATURE:
<223> OTHER INFORMATION: Mouse polylysine
<400> SEQUENCE: 2

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Lys Lys Lys Arg Arg Lys Lys Lys Lys Gly Lys Gly Ala Val Ser Ala
 1               5                  10                  15
Val Gln Gln Glu Leu Asp Lys Glu Ser Gly Ala Leu Val Asp Glu Val
            20                  25                  30
Ala Lys Gln Leu Glu Ser Gln Ala Leu Glu Glu Lys Glu Arg Asp Asp
        35                  40                  45
Asp Asp Glu Asp Gly Asp Gly Asp Ala Asp Gly Ala Thr Gly Lys Lys
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66      50                  55                  60
68 Lys Lys Lys Lys Lys Lys Lys
69  65                  70
72 <210> SEQ ID NO: 3
73 <211> LENGTH: 57
74 <212> TYPE: PRT
75 <213> ORGANISM: Saccharomyces sp.
77 <220> FEATURE:
78 <223> OTHER INFORMATION: Saccharomyces polylysine
80 <400> SEQUENCE: 3
81 Thr Asp Ala Glu Ile Glu Asn Ser Pro Ala Ser Asp Leu Lys Glu Leu
82   1               5                  10                  15
84 Asn Leu Glu Asn Glu Gly Val Glu Gln Gln Asp Gln Ala Lys Ala Asp
85              20                  25                  30
87 Glu Ser Asp Pro Val Glu Ser Lys Lys Lys Lys Asn Lys Lys Lys Lys
88          35                  40                  45
90 Lys Lys Lys Ser Asn Val Lys Lys Ile
91      50                  55
94 <210> SEQ ID NO: 4
95 <211> LENGTH: 35
96 <212> TYPE: DNA
97 <213> ORGANISM: Artificial Sequence
99 <220> FEATURE:
100 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
101       oligonucleotide
103 <400> SEQUENCE: 4
104 caaccattgt gctgcagctt tcacacccaa tgcag                              35
107 <210> SEQ ID NO: 5
108 <211> LENGTH: 35
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:
113 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
114       oligonucleotide
116 <400> SEQUENCE: 5
117 ctgcattggg tgtgaaagct gcagcacaat ggttg                              35
120 <210> SEQ ID NO: 6
121 <211> LENGTH: 478
122 <212> TYPE: PRT
123 <213> ORGANISM: Homo Sapiens
125 <220> FEATURE:
126 <223> OTHER INFORMATION: Human dnvMetAP2
128 <220> FEATURE:
129 <221> NAME/KEY: SITE
130 <222> LOCATION: (219)
131 <223> OTHER INFORMATION: May be any naturally occurring amino acid
133 <220> FEATURE:
134 <221> NAME/KEY: SITE
135 <222> LOCATION: (231)
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RAW SEQUENCE LISTING DATE: 10/25/2004
PATENT APPLICATION: **US/10/712,359A** TIME: 11:45:50

Input Set : **A:\27908007.app**
Output Set: **N:\CRF4\10252004\J712359A.raw**

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136 <223> OTHER INFORMATION: May be any amino acid, except His
138 <220> FEATURE:
139 <221> NAME/KEY: SITE
140 <222> LOCATION: (251)
141 <223> OTHER INFORMATION: May be any naturally occurring amino acid
143 <220> FEATURE:
144 <221> NAME/KEY: SITE
145 <222> LOCATION: (262)
146 <223> OTHER INFORMATION: May be any naturally occurring amino acid
148 <220> FEATURE:
149 <221> NAME/KEY: SITE
150 <222> LOCATION: (328)
151 <223> OTHER INFORMATION: May be any naturally occurring amino acid
153 <220> FEATURE:
154 <221> NAME/KEY: SITE
155 <222> LOCATION: (331)
156 <223> OTHER INFORMATION: May be any naturally occurring amino acid
158 <220> FEATURE:
159 <221> NAME/KEY: SITE
160 <222> LOCATION: (338)..(339)
161 <223> OTHER INFORMATION: May be any naturally occurring amino acid
163 <220> FEATURE:
164 <221> NAME/KEY: SITE
165 <222> LOCATION: (364)
166 <223> OTHER INFORMATION: May be any naturally occurring amino acid
168 <220> FEATURE:
169 <221> NAME/KEY: SITE
170 <222> LOCATION: (444)
171 <223> OTHER INFORMATION: May be any naturally occurring amino acid
173 <220> FEATURE:
174 <221> NAME/KEY: SITE
175 <222> LOCATION: (447)
176 <223> OTHER INFORMATION: May be any naturally occurring amino acid
178 <220> FEATURE:
179 <221> NAME/KEY: SITE
180 <222> LOCATION: (459)
181 <223> OTHER INFORMATION: May be any naturally occurring amino acid
183 <400> SEQUENCE: 6
184 Met Ala Gly Val Glu Glu Val Ala Ala Ser Gly Ser His Leu Asn Gly
185   1               5                  10                  15
187 Asp Leu Asp Pro Asp Asp Arg Glu Glu Gly Ala Ala Ser Thr Ala Glu
188              20                  25                  30
190 Glu Ala Ala Lys Lys Lys Arg Arg Lys Lys Lys Lys Ser Lys Gly Pro
191          35                  40                  45
193 Ser Ala Ala Gly Glu Gln Glu Pro Asp Lys Glu Ser Gly Ala Ser Val
194      50                  55                  60
196 Asp Glu Val Ala Arg Gln Leu Glu Arg Ser Ala Leu Glu Asp Lys Glu
197  65                  70                  75                  80
199 Arg Asp Glu Asp Asp Glu Asp Gly Asp Gly Asp Gly Asp Gly Ala Thr
```

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Input Set : **A:\27908007.app**
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```
200                 85                  90                  95
202 Gly Lys Lys Lys Lys Lys Lys Lys Lys Arg Gly Pro Lys Val Gln
203             100                 105                 110
205 Thr Asp Pro Pro Ser Val Pro Ile Cys Asp Leu Tyr Pro Asn Gly Val
206         115                 120                 125
208 Phe Pro Lys Gly Gln Glu Cys Glu Tyr Pro Pro Thr Gln Asp Gly Arg
209     130                 135                 140
211 Thr Ala Ala Trp Arg Thr Thr Ser Glu Glu Lys Lys Ala Leu Asp Gln
212 145                 150                 155                 160
214 Ala Ser Glu Glu Ile Trp Asn Asp Phe Arg Glu Ala Ala Glu Ala His
215                 165                 170                 175
217 Arg Gln Val Arg Lys Tyr Val Met Ser Trp Ile Lys Pro Gly Met Thr
218             180                 185                 190
220 Met Ile Glu Ile Cys Glu Lys Leu Glu Asp Cys Ser Arg Lys Leu Ile
221         195                 200                 205
W--> 223 Lys Glu Asn Gly Leu Asn Ala Gly Leu Ala Xaa Pro Thr Gly Cys Ser
224     210                 215                 220
226 Leu Asn Asn Cys Ala Ala Xaa Tyr Thr Pro Asn Ala Gly Asp Thr Thr
227 225                 230                 235                 240
229 Val Leu Gln Tyr Asp Asp Ile Cys Lys Ile Xaa Phe Gly Thr His Ile
230                 245                 250                 255
232 Ser Gly Arg Ile Ile Xaa Cys Ala Phe Thr Val Thr Phe Asn Pro Lys
233             260                 265                 270
235 Tyr Asp Thr Leu Leu Lys Ala Val Lys Asp Ala Thr Asn Thr Gly Ile
236         275                 280                 285
238 Lys Cys Ala Gly Ile Asp Val Arg Leu Cys Asp Val Gly Glu Ala Ile
239     290                 295                 300
241 Gln Glu Val Met Glu Ser Tyr Glu Val Glu Ile Asp Gly Lys Thr Tyr
242 305                 310                 315                 320
244 Gln Val Lys Pro Ile Arg Asn Xaa Asn Gly Xaa Ser Ile Gly Gln Tyr
245                 325                 330                 335
247 Arg Xaa Xaa Ala Gly Lys Thr Val Pro Ile Val Lys Gly Gly Glu Ala
248             340                 345                 350
250 Thr Arg Met Glu Glu Gly Glu Val Tyr Ala Ile Xaa Thr Phe Gly Ser
251         355                 360                 365
253 Thr Gly Lys Gly Val Val His Asp Asp Met Glu Cys Ser His Tyr Met
254     370                 375                 380
256 Lys Asn Phe Asp Val Gly His Val Pro Ile Arg Leu Pro Arg Thr Lys
257 385                 390                 395                 400
259 His Leu Leu Asn Val Ile Asn Glu Asn Phe Gly Thr Leu Ala Phe Cys
260                 405                 410                 415
262 Arg Arg Trp Leu Asp Arg Leu Gly Glu Ser Lys Tyr Leu Met Ala Leu
263             420                 425                 430
265 Lys Asn Leu Cys Asp Leu Gly Ile Val Asp Pro Xaa Pro Pro Xaa Cys
266         435                 440                 445
268 Asp Ile Lys Gly Ser Tyr Thr Ala Gln Phe Xaa His Thr Ile Leu Leu
269     450                 455                 460
271 Arg Pro Thr Cys Lys Glu Val Val Ser Arg Gly Asp Asp Tyr
272 465                 470                 475
```

RAW SEQUENCE LISTING DATE: 10/25/2004
PATENT APPLICATION: **US/10/712,359A** TIME: 11:45:50

Input Set : **A:\27908007.app**
Output Set: **N:\CRF4\10252004\J712359A.raw**

275 <210> SEQ ID NO: 7
276 <211> LENGTH: 478
277 <212> TYPE: PRT
278 <213> ORGANISM: Mus musculus
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Mouse MetAP2
283 <220> FEATURE:
284 <221> NAME/KEY: SITE
285 <222> LOCATION: (219)
286 <223> OTHER INFORMATION: May be any naturally occurring amino acid
288 <220> FEATURE:
289 <221> NAME/KEY: SITE
290 <222> LOCATION: (231)
291 <223> OTHER INFORMATION: May be any amino acid, except His
293 <220> FEATURE:
294 <221> NAME/KEY: SITE
295 <222> LOCATION: (251)
296 <223> OTHER INFORMATION: May be any naturally occurring amino acid
298 <220> FEATURE:
299 <221> NAME/KEY: SITE
300 <222> LOCATION: (262)
301 <223> OTHER INFORMATION: May be any naturally occurring amino acid
303 <220> FEATURE:
304 <221> NAME/KEY: SITE
305 <222> LOCATION: (328)
306 <223> OTHER INFORMATION: May be any naturally occurring amino acid
308 <220> FEATURE:
309 <221> NAME/KEY: SITE
310 <222> LOCATION: (331)
311 <223> OTHER INFORMATION: May be any naturally occurring amino acid
313 <220> FEATURE:
314 <221> NAME/KEY: SITE
315 <222> LOCATION: (338)..(339)
316 <223> OTHER INFORMATION: May be any naturally occurring amino acid
318 <220> FEATURE:
319 <221> NAME/KEY: SITE
320 <222> LOCATION: (364)
321 <223> OTHER INFORMATION: May be any naturally occurring amino acid
323 <220> FEATURE:
324 <221> NAME/KEY: SITE
325 <222> LOCATION: (444)
326 <223> OTHER INFORMATION: May be any naturally occurring amino acid
328 <220> FEATURE:
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330 <222> LOCATION: (447)
331 <223> OTHER INFORMATION: May be any naturally occurring amino acid
333 <220> FEATURE:
334 <221> NAME/KEY: SITE
335 <222> LOCATION: (459)

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 10/25/2004
PATENT APPLICATION: **US/10/712,359A** TIME: 11:45:51

Input Set : **A:\27908007.app**
Output Set: **N:\CRF4\10252004\J712359A.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; Xaa Pos. 219,231,251,262,328,331,338,339,364,444,447,459
Seq#:7; Xaa Pos. 219,231,251,262,328,331,338,339,364,444,447,459
Seq#:8; Xaa Pos. 162,174,194,205,271,274,281,282,307,387,390,402
Seq#:9; N Pos. 693
Seq#:10; N Pos. 693
Seq#:11; N Pos. 522
Seq#:16; Xaa Pos. 219,231,251,262,328,331,338,339,364,444,447,459
Seq#:18; N Pos. 779

VERIFICATION SUMMARY DATE: 10/25/2004
PATENT APPLICATION: **US/10/712,359A** TIME: 11:45:51

Input Set : **A:\27908007.app**
Output Set: **N:\CRF4\10252004\J712359A.raw**

L:223 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:208
M:341 Repeated in SeqNo=6
L:378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:208
M:341 Repeated in SeqNo=7
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:160
M:341 Repeated in SeqNo=8
L:601 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:660
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:660
L:678 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:480
L:1116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:208
M:341 Repeated in SeqNo=16
L:1294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:720